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PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/09/757,673

DATE: 03/14/2001 TIME: 04:18:43

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This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING ENTERED
2		
3 4	(1) G	eneral Information:
5 6 7	(i)	APPLICANT: Wilson, James M. Fisher, Krishna J.
, 8 9 10	(ii)	TITLE OF INVENTION: Method for Recombinant Adeno-Associated Virus-Directed Gene Therapy
11 12	(iii)	NUMBER OF SEQUENCES: 4
13 14 15 16 17 18 19 20	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Howson and Howson (B) STREET: Spring House Corporate Cntr, PO Box 457 (C) CITY: Spring House (D) STATE: Pennsylvania (E) COUNTRY: USA (F) ZIP: 19477
21 22 23 24 25 26	(v)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
27 28 29 30 31	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: 09/757,673 (B) FILING DATE: (C) CLASSIFICATION:
32 33 34 35	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 09/242,977 (B) FILING DATE:
36 37 38 39	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: US 08/729,061 (B) FILING DATE: 10-OCT-1996
40 41	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: Kodroff, Cathy A.
42 43 44		(B) REGISTRATION NUMBER: 33,980 (C) REFERENCE/DOCKET NUMBER: GNVPN.019CIP2USA
45 46	(ix)	TELECOMMUNICATION INFORMATION: (A) TELEPHONE: 215-540-9200

RAW SEQUENCE LISTING PATENT APPLICATION US/09/757,673

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47	(B) TELEFAX: 215-540-5818			
48				
49				
50	(2)			
51	(2) INFORMATION FOR SEQ ID NO:1:			
52	(;) GEOLIENGE CHARACHERICHICG.			
53 54	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10398 base pairs			
55	(B) TYPE: nucleic acid			
56	(C) STRANDEDNESS: double			
57	(D) TOPOLOGY: unknown			
58	(-,			
59	(ii) MOLECULE TYPE: cDNA			
60				
61				
62				
63	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:			
64				
65	GAATTCGCTA GCATCATCAA TAATATACCT TATTTTGGAT TGAAGCCAA	r Argaraarga	60	
66 67	GGGGGTGGAG TTTGTGACGT GGCGCGGGGC GTGGGAACGG GGCGGGTGA	ር ርጥአርጥአርጥርጥ	120	
68	GGGGGTGGAG TITGTGACGT GGCGCGGGC GTGGGAACGG GGCGGGTGA	C GIAGIAGIGI	120	
69	GGCGGAAGTG TGATGTTGCA AGTGTGGCGG AACACATGTA AGCGACGGA	r grggcaaaag	180	
70				
71	TGACGTTTTT GGTGTGCGCC GGTGTACACA GGAAGTGACA ATTTTCGCG	C GGTTTTAGGC	240	
72	•			
73	GGATGTTGTA GTAAATTTGG GCGTAACCGA GTAAGATTTG GCCATTTTC	G CGGGAAAACT	300	
74				
75	GAATAAGAGG AAGTGAAATC TGAATAATTT TGTGTTACTC ATAGCGCGT.	A ATATTTGTCT	360	
76	ACCORANTED COMESCOCION COCHEGOCIONA CINCACCOCCO COCCCONA NA	а даааааааа	400	
77 78	AGGGAGATCT GCTGCGCGCT CGCTCGCTCA CTGAGGCCGC CCGGGCAAA	3 CCCGGGCGTC	420	
79	GGGCGACCTT TGGTCGCCCG GCCTCAGTGA GCGAGCGAGC GCGCAGAGA	с ссастсссса	480	
80	GGGGGGCTT TGGTCGCCG GCCTCAGTGA GCGAGCGAGC GCGCAGAGA	J GOAGTOGECA	100	
81	ACTCCATCAC TAGGGGTTCC TTGTAGTTAA TGATTAACCC GCCATGCTA	C TTATCTACAA	540	
82				
83	TTCGAGCTTG CATGCCTGCA GGTCGTTACA TAACTTACGG TAAATGGCCC	C GCCTGGCTGA	600	
84				
85	CCGCCCAACG ACCCCCGCCC ATTGACGTCA ATAATGACGT ATGTTCCCA	r agtaacgcca	660	
86				
87	ATAGGGACTT TCCATTGACG TCAATGGGTG GAGTATTTAC GGTAAACTG	C CCACTTGGCA	720	
88 89	GTACATCAAG TGTATCATAT GCCAAGTACG CCCCCTATTG ACGTCAATGA	\	780	
90	GIACATCAAG IGIATCATAT GCCAAGTACG CCCCCTATTG ACGTCAATGA	COGIAAAIGG	700	
91	CCCGCCTGGC ATTATGCCCA GTACATGACC TTATGGGACT TTCCTACTTC	GCAGTACATC :	840	
92			-	
93	TACGTATTAG TCATCGCTAT TACCATGGTG ATGCGGTTTT GGCAGTACA	CAATGGGCGT	900	
94				
95	GGATAGCGGT TTGACTCACG GGGATTTCCA AGTCTCCACC CCATTGACG	CAATGGGAGT	960	
96				
97	TTGTTTTGGC ACCAAAATCA ACGGGACTTT CCAAAATGTC GTAACAACTC	C CGCCCCATTG 10	020	
98				

ACGCAAATGG GCGGTAGGCG TGTACGGTGG GAGGTCTATA TAAGCAGAGC TCGTTTAGTG

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					-	IVI OI DELI. DOU	2071.417
100 101	AACCGTCAGA	TCGCCTGGAG	ACGCCATCCA	CGCTGTTTTG	ACCTCCATAG	AAGACACCGG	1140
102							
103 104	GACCGATCCA	GCCTCCGGAC	TCTAGAGGAT	CCGGTACTCG	AGGAACTGAA	AAACCAGAAA	1200
105	GTTAACTGGT	AAGTTTAGTC	TTTTTGTCTT	TTATTTCAGG	TCCCGGATCC	GGTGGTGGTG	1260
106	CAAATCAAAG	AACTGCTCCT	CAGTGGATGT	TGCCTTTACT	TCTAGGCCTG	TACGGAAGTG	1320
108	TTACTTCTGC	TCTAAAAGCT	GCGGAATTGT	ACCCGCGGCC	GCAATTCCCG	GGGATCGAAA	1380
110	GAGCCTGCTA	AAGCAAAAA	GAAGTCACCA	TGTCGTTTAC	TTTGACCAAC	AAGAACGTGA	1440
112			~~~~ 	maar ar aar a	G3 3 GG3 GGEG	CEC TO COCO	7.500
113 114	TTTTCGTTGC	CGGTCTGGGA	GGCATTGGTC	TGGACACCAG	CAAGGAGCTG	CTCAAGCGCG	1500
115 116	ATCCCGTCGT	TTTACAACGT	CGTGACTGGG	AAAACCCTGG	CGTTACCCAA	CTTAATCGCC	1560
117 118	TTGCAGCACA	TCCCCCTTTC	GCCAGCTGGC	GTAATAGCGA	AGAGGCCCGC	ACCGATCGCC	1620
119 120	CTTCCCAACA	GTTGCGCAGC	CTGAATGGCG	AATGGCGCTT	TGCCTGGTTT	CCGGCACCAG	1680
121	AAGCGGTGCC	GGAAAGCTGG	CTGGAGTGCG	ATCTTCCTGA	GGCCGATACT	GTCGTCGTCC	1740
122	CCTCAAACTG	GCAGATGCAC	GGTTACGATG	CGCCCATCTA	CACCAACGTA	ACCTATCCCA	1800
124 125	TTACGGTCAA	TCCGCCGTTT	GTTCCCACGG	AGAATCCGAC	GGGTTGTTAC	TCGCTCACAT	1860
126 127	TTAATGTTGA	TGAAAGCTGG	CTACAGGAAG	GCCAGACGCG	AATTATTTTT	GATGGCGTTA	1920
128 129	ACTCGGCGTT	TCATCTGTGG	TGCAACGGGC	GCTGGGTCGG	TTACGGCCAG	GACAGTCGTT	1980
130 131	TGCCGTCTGA	ATTTGACCTG	AGCGCATTTT	TACGCGCCGG	AGAAAACCGC	CTCGCGGTGA	2040
132							
133 134	TGGTGCTGCG	TTGGAGTGAC	GGCAGTTATC	TGGAAGATCA	GGATATGTGG	CGGATGAGCG	2100
135 136	GCATTTTCCG	TGACGTCTCG	TTGCTGCATA	AACCGACTAC	ACAAATCAGC	GATTTCCATG	2160
137 138	TTGCCACTCG	CTTTAATGAT	GATTTCAGCC	GCGCTGTACT	GGAGGCTGAA	GTTCAGATGT	2220
139 140	GCGGCGAGTT	GCGTGACTAC	CTACGGGTAA	CAGTTTCTTT	ATGGCAGGGT	GAAACGCAGG	2280
141 142	TCGCCAGCGG	CACCGCGCCT	TTCGGCGGTG	AAATTATCGA	TGAGCGTGGT	GGTTATGCCG	2340
143	ATCGCGTCAC	ACTACGTCTG	AACGTCGAAA	ACCCGAAACT	GTGGAGCGCC	GAAATCCCGA	2400
144 145	ATCTCTATCG	TGCGGTGGTT	GAACTGCACA	CCGCCGACGG	CACGCTGATT	GAAGCAGAAG	2460
146 147	CCTGCGATGT	CGGTTTCCGC	GAGGTGCGGA	TTGAAAATGG	TCTGCTGCTG	CTGAACGGCA	2520
148 149	AGCCGTTGCT	GATTCGAGGC	GTTAACCGTC	ACGAGCATCA	TCCTCTGCAT	GGTCAGGTCA	2580
150 151 152	TGGATGAGCA	GACGATGGTG	CAGGATATCC	TGCTGATGAA	GCAGAACAAC	TTTAACGCCG	2640

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153 154	TGCGCTGTTC	GCATTATCCG	AACCATCCGC	TGTGGTACAC	_		2700
155 156	TGTATGTGGT	GGATGAAGCC	AATATTGAAA	CCCACGGCAT	GGTGCCAATG	AATCGTCTGA	2760
157 158	CCGATGATCC	GCGCTGGCTA	CCGGCGATGA	GCGAACGCGT	AACGCGAATG	GTGCAGCGCG	2820
159 160	ATCGTAATCA	CCCGAGTGTG	ATCATCTGGT	CGCTGGGGAA	TGAATCAGGC	CACGGCGCTA	2880
161 162	ATCACGACGC	GCTGTATCGC	TGGATCAAAT	CTGTCGATCC	TTCCCGCCCG	GTGCAGTATG	2940
163 164	AAGGCGGCGG	AGCCGACACC	ACGGCCACCG	ATATTATTTG	CCCGATGTAC	GCGCGCGTGG	3000
165 166	ATGAAGACCA	GCCCTTCCCG	GCTGTGCCGA	AATGGTCCAT	CAAAAAATGG	CTTTCGCTAC	3060
167 168	CTGGAGAGAC	GCGCCCGCTG	ATCCTTTGCG	AATACGCCCA	CGCGATGGGT	AACAGTCTTG	3120
169 170			,	GTCAGTATCC			3180
171 172				AATATGATGA			3240
173 174				ACGATCGCCA			3300
175 176	_			TGACGGAAGC			3360
177 178				AAGTGACCAG			3420
179 180				CGCTGGATGG		1	3480
181 182				AACAGTTGAT CAGTACGCGT			3540 3600
183 184 185			•	GGCAGCAGTG		•	3660
186 187				TCCCGCATCT			3720
188 189				AATTTAACCG			3780
190 191				TGACGCCGCT			3840
192 193				AAGCGACCCG			3900
194 195						CAGTGCACGG	3960
196 197	CAGATACACT	TGCTGATGCG	GTGCTGATTA	CGACCGCTCA	CGCGTGGCAG	CATCAGGGGA	4020
198 199	AAACCTTATT	TATCAGCCGG	AAAACCTACC	GGATTGATGG	TAGTGGTCAA	ATGGCGATTA	4080
200 201	CCGTTGATGT	TGAAGTGGCG	AGCGATACAC	CGCATCCGGC	GCGGATTGGC	CTGAACTGCC	4140
202 203	AGCTGGCGCA	GGTAGCAGAG	CGGGTAAACT	GGCTCGGATT	AGGGCCGCAA	GAAAACTATC	4200
204 205	CCGACCGCCT	TACTGCCGCC	TGTTTTGACC	GCTGGGATCT	GCCATTGTCA	GACATGTATA	4260

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208	00000111001	CIICCOAGC	. GAMMACGGIC	. 100001000	GACGCGCGAA	TIGAATTATG	4320
209	GCCCACACC	СТСССССССС	י מאפייירים אפיי	י ייירא א ראיירא רי	CCCCTT C C CT	' CAACAGCAAC	4200
210	00001011001	. 0100000000	. OACTICCAGE	. ICAACAICAG	CCGCIACAGI	CAACAGCAAC	4380
211	ТСАТССААА С	CACCCATCCC	י מאיירייים מייים	י אכככככאאכא	3.C.C.C.3.C.3.E.C.C	CTGAATATCG	4440
212	IGNIGGAMAC	CAGCCATCGC	. CAICIGCIGC	. ACGCGGAAGA	AGGCACATGG	CTGAATATCG	4440
212	A CCCTTTTCCA		COMOGOO A GO	1 2 CMCCMCC2 C	GGGGTG3 GT-		
213	ACGGIIICCA	IAIGGGGAII	GGTGGCGACG	ACTCCTGGAG	CCCGTCAGTA	TCGGCGGAAT	4500
	ma da domoa o						
215	TACAGCTGAG	CGCCGGTCGC	TACCATTACC	: AGTTGGTCTG	GTGTCAAAAA	TAATAATAAC	4560
216	~~~~~						
217	CGGGCAGGCC	ATGTCTGCCC	GTATTTCGCG	TAAGGAAATC	CATTATGTAC	TATTTAAAAA	4620
218							
219	ACACAAACTT	' TTGGATGTTC	GGTTTATTCT	TTTTCTTTTA	CTTTTTTATC	ATGGGAGCCT	4680
220							
221	ACTTCCCGTT	TTTCCCGATT	TGGCTACATG	ACATCAACCA	TATCAGCAAA	AGTGATACGG	4740
222							
223	GTATTATTTT	TGCCGCTATT	TCTCTGTTCT	CGCTATTATT	CCAACCGCTG	TTTGGTCTGC	4800
224							
225	TTTCTGACAA	ACTCGGCCTC	GACTCTAGGC	GGCCGCGGG	ATCCAGACAT	GATAAGATAC	4860
226						0	1000
227	ATTGATGAGT	TTGGACAAAC	CACAACTAGA	ATGCAGTGAA	ΔΔΔΔΤΩΟΤΤ	TATTTGTGAA	4920
228						INITIOIONA	4920
229	ATTTGTGATG	СФАТТССТТТ	ል ጥጥርጥልልሮር	ATTATAAGCT	CCAATAAACA	አርጣሞን አርን አር	4000
230		01111100111	ATTIOTACC	ATTALAAGCI	GCAATAAACA	AGITAACAAC	4980
231	አ ልሮልልሞሞርሮል	ጥጥር እ ጥጥጥጥ አጥ	CTTTTC A CCTTT	CAGGGGGAGG	-	MMMMM CCC 2 m	
232	MACAMITOCA	IICAIIIIAI	GITICAGGIT	CAGGGGGAGG	TGTGGGAGGT	TTTTTCGGAT	5040
232	CCTCTACACT	CCACMACAMA	3 C	CCCCCCC A A TCC			
234	CCICIAGAGI	CGAGTAGATA	AGTAGCATGG	CGGGTTAATC	ATTAACTACA	AGGAACCCCT	5100
235	3.CMC3.MCC3.C	mmaaaaa ama				,	
	AGIGAIGGAG	TTGGCCACTC	CCTCTCTGCG	CGCTCGCTCG	CTCACTGAGG	CCGGGCGACC	5160
236	11166maaa	227 222 222	*				
237	AAAGGTCGCC	CGACGCCCGG	GCTTTGCCCG	GGCGGCCTCA	GTGAGCGAGC	GAGCGCGCAG	5220
238							
239	CAGATCTGGA	AGGTGCTGAG	GTACGATGAG	ACCCGCACCA	GGTGCAGACC	CTGCGAGTGT	5280
240							
241	GGCGGTAAAC	ATATTAGGAA	CCAGCCTGTG	ATGCTGGATG	TGACCGAGGA	GCTGAGGCCC	5340
242							
243	GATCACTTGG	TGCTGGCCTG	CACCCGCGCT	GAGTTTGGCT	CTAGCGATGA	AGATACAGAT	5400
244							
245	TGAGGTACTG	AAATGTGTGG	GCGTGGCTTA	AGGGTGGGAA	AGAATATATA	AGGTGGGGGT	5460
246							
247	CTTATGTAGT	TTTGTATCTG	TTTTGCAGCA	GCCGCCGCCG	CCATGAGCAC	CAACTCGTTT	5520
248						0.1.0100111	3320
249	GATGGAAGCA	TTGTGAGCTC	ATATTTGACA	ACGCGCATGC	CCCCATGGGC	СССССТСССТ	5580
250					00007110000	0000010001	2200
251	CAGAATGTGA	TGGGCTCC≱G	САТТСАТССТ	רפרררתיירי	ጥር ር ር ር ር ር ሊ እ እ	CTCTACTACC	5640
252		- JOSSI COMO		2322223100	1 GCCCGCAAA	CICIACIACC	2040
253	ТТСАССТАСС	ΔGΔCCCTCTC	TCCAACCCC	TTGGAGACTG	CVCCCMCCC	CCCCCCCCCCCCC	E700
254	-10		1 JOANCOCCG	TIGGAGACIG	CAGCCTCCGC	CGCCGCTTCA	5700
255	GCCGCTCCAC	CCACCCCCC	CCCCAmmama	ACTGACTTTG	ammaama ~	aaaaammaa.	
256	CCCCTGCAG	CCACCGCCCG	CGGGAIIGIG	ACTGACTTTG	CTTTCCTGAG	CCCGCTTGCA	5760
257	ACCACTCCAC	CTTTCCCCTTTC	ATTCCCCCCCCC	G3.0003.033.00	mas aac		
258	AGCAGIGCAG	CITCCCGITC	ATCCGCCCGC	GATGACAAGT	TGACGGCTCT	TTTGGCACAA	5820
250							

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